

DESCRIPTION

REGULATOR FOR ADIPONECTIN RECEPTOR EXPRESSION

5 Technical Field

The present invention relates to therapies for insulin resistance and diabetes that utilize regulation of adiponectin receptor expression.

Background Art

10 Adiponectin/Acrp30 (Non-Patent Documents 1-4) is a hormone secreted from adipocytes, which acts as an anti-diabetic (Non-Patent Documents 5-12) and anti-atherogenic (Non-Patent Documents 8, 12, and 13) adipokine. Adiponectin is thought to accentuate insulin-sensitivity by enhancing fatty acid oxidation via the activation of 5'-AMP-activated protein kinase (AMPK) (Non-Patent Documents 10 and
15 11) and peroxisome proliferator-activated receptor (PPAR)- α (Non-Patent Documents 5, 6, and 12). Recently, the present inventors succeeded in the cloning of cDNAs encoding adiponectin receptors 1 and 2 (AdipoR1, and AdipoR2) (Patent Document 1, Non-Patent Document 14). AdipoR1 is abundantly expressed in skeletal muscle, whereas AdipoR2 is primarily expressed in liver. AdipoR1 and R2 are predicted to
20 contain seven transmembrane domains (Non-Patent Document 14), but are considered to be structurally and functionally distinct from G-protein-coupled receptors (Non-Patent Documents 15-17). AdipoR1 and R2 function as receptors for globular adiponectin and full-length adiponectin, and mediate the enhancement of AMPK activity (Non-Patent Documents 10 and 11); PPAR α ligand activity (Non-Patent
25 Document 12), and fatty acid oxidation and glucose uptake by adiponectin (Non-Patent Document 14).

As described above, findings on AdipoR1 and AdipoR2 are starting to accumulate, but it was not known whether the expression of AdipoR1 and R2 changes under physiological and pathophysiological conditions.

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[Patent Document 1] International Application Number: PCT/JP2003/007515,
International Application Number: WO2004/061108, Patent number: US2004241802
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Disclosure of the Invention

The first objective of the present invention is to elucidate the physiological and pathophysiological functions of adiponectin receptors (AdipoR1 and AdipoR2). The second objective is to find a means for applying adiponectin receptors in diabetes treatment, based on the above functions.

In order to solve the above issues, the present inventors examined the expression levels of AdipoR1 and AdipoR2 and the insulin levels in mice under fasting and refeeding conditions. The AdipoR1/R2 expressions dramatically changed by fasting and refeeding. Such changes in AdipoR1/R2 expressions were suggested to be inversely correlated with *in vivo* plasma insulin levels, indicating that AdipoR1/R2 may be important regulators of the metabolic sensitivity to nutritive conditions and insulin.

Next, the AdipoR1/R2 expressions were analyzed in a mouse model in which insulin deficiency was induced by streptozotocin (STZ) treatment (Non-Patent Document 18). The STZ treatment enhanced the AdipoR1/R2 mRNA levels significantly in the liver, skeletal muscle, and adipose tissues of mice, which was shown to be restored by insulin administration. Moreover, the reduction of AdipoR1/R2 mRNAs in myocytes and hepatocytes was confirmed *in vitro*.

Analyses of the AdipoR1/R2 expressions in a mouse model of hyperinsulinemia caused by obesity-related insulin resistance (ob/ob mice) showed that expressions of AdipoR1/R2 were significantly reduced in the muscles and adipose tissues of ob/ob mice. The binding of adiponectin to membrane fractions of the above tissues was significantly decreased in ob/ob mice as compared to wild type mice, and AMP kinase activation by adiponectin was also decreased in the above tissues of ob/ob mice.

The above results reveal that insulin negatively regulates the expression levels of adiponectin receptors and adiponectin sensitivity, and that obesity-related insulin resistance/hyperinsulinemia causes downregulation of adiponectin receptors and adiponectin resistance.

Furthermore, the present inventors analyzed the mechanism behind insulin-mediated decrease of AdipoR1/R2 gene expression. Interestingly, the PI3-kinase inhibitor LY294002 negates the AdipoR1/R2 mRNA level reduction effect of insulin, whereas PD98059, a mitogen-activated protein kinase activation inhibitor, did not affect the above effect of insulin. Thus, the downregulation of adiponectin receptors by insulin was suggested to be mediated by the PI3-kinase signaling pathway.

Moreover, the present inventors examined the events downstream of PI3-kinase regarding the regulation of adiponectin receptor expression by insulin. It has been found that Akt, a downstream effector of PI3-kinase, phosphorylates transcriptional activators of the forkhead family, which results in inhibition of transcriptional activities of these phosphorylated transcriptional activators. On the other hand, AdipoR1/R2 genes comprise in their promoter an element capable of binding to the forkhead family transcriptional activator, FoxoI. Therefore, mouse FoxoI (wild type and constitutively active type) was transfected into the C2C12 cell line, and then the cell line was treated with insulin to observe AdipoR1/R2 expressions. The results showed that FoxoI enhanced AdipoR1/R2 expressions, and that insulin downregulates adiponectin receptor expression via inactivation of FoxoI.

In summary, the present inventors revealed, for the first time, that AdipoR1/R2 expressions in insulin target tissues are inversely correlated with plasma insulin levels, and that insulin negatively regulates the expression levels of adiponectin receptors through the PI3-kinase/FoxoI pathway. Not only AdipoR1/R2 agonists, but also a strategy to increase AdipoR1/R2 was shown to be effective for the first time, for the treatment of insulin resistance and type 2 diabetes. Thus, the present invention relates to therapeutic agents for diabetes mediating AdipoR1/R2 regulation, and more specifically provides the following:

- [1] An adiponectin receptor expression-enhancing agent, comprising a FoxoI protein or a FoxoI gene.
- [2] An adiponectin resistance-improving agent, comprising a FoxoI protein or a FoxoI gene as an active ingredient.
- [3] An insulin resistance-improving agent, comprising a FoxoI protein or a FoxoI gene as an active ingredient.

- [4] An obesity-preventing or improving agent, comprising a FoxoI protein or a FoxoI gene as an active ingredient.
- [5] A diabetes-preventing or treating agent, comprising a FoxoI protein or a FoxoI gene as an active ingredient.
- 5 [6] An arteriosclerosis-preventing or treating agent, comprising a FoxoI protein or a FoxoI gene as an active ingredient.
- [7] An adiponectin receptor expression-enhancing agent, comprising a PI3 (Phosphoinositide 3)-kinase-FoxoI pathway inhibitor.
- [8] The adiponectin receptor expression-enhancing agent of [7], wherein the
- 10 [9] The adiponectin receptor expression-enhancing agent of [8], wherein the PI3-kinase inhibitor is LY294002.
- [10] An adiponectin resistance-improving agent, comprising as an active ingredient the adiponectin receptor expression-enhancing agent of any one of [7] to [9].
- 15 [11] An insulin resistance-improving agent, comprising as an active ingredient the adiponectin receptor expression-enhancing agent of any one of [7] to [9].
- [12] An obesity-preventing or improving agent, comprising as an active ingredient the adiponectin receptor expression-enhancing agent of any one of [7] to [9].
- [13] A diabetes-preventing or treating agent, comprising as an active ingredient the
- 20 [14] An arteriosclerosis-preventing or treating agent, comprising as an active ingredient the adiponectin receptor expression-enhancing agent of any one of [7] to [9].
- [15] An adiponectin receptor expression-suppressing agent, comprising insulin or insulin gene.
- 25 [16] A method of screening for an adiponectin receptor expression-enhancing agent, an adiponectin resistance-improving agent, an insulin resistance-improving agent, an obesity-improving agent, a therapeutic agent for diabetes, or a therapeutic agent for arteriosclerosis, comprising the step of contacting a test substance with PI3-kinase and a PI3-kinase substrate.
- 30 [17] A method of screening for an adiponectin receptor expression-enhancing agent, an adiponectin resistance-improving agent, an insulin resistance-improving agent, an obesity-improving agent, a therapeutic agent for diabetes, or a therapeutic agent for arteriosclerosis, wherein the method comprises the step of contacting a test substance with Akt and FoxoI.

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The present invention revealed that insulin is involved in the regulation of

adiponectin receptors and the mechanism of regulation, and at the same time, achieved specific agents for enhancing adiponectin receptor expression for the first time.

Agents that enhance the expression of adiponectin receptors resolve both adiponectin resistance and insulin resistance, and thus, will be useful as novel agents for treating

5 insulin resistance or type 2 diabetes.

Brief Description of the Drawings

Fig. 1 shows the effects of nutrition control on the expression of AdipoR1/R2 genes. The expression levels of AdipoR1 (A and C) and AdipoR2 (B and D) in liver (A and B) and skeletal muscle (A and B), the blood glucose level (E), and the plasma insulin level (F) of mice were compared under conditions of unrestricted feeding, fasting, and refeeding. The unrestricted feeding group (ad lib) could feed freely (lane 1). The fasted group was separated from food for 48 hrs (lane 2). The refed group was allowed to freely feed for 6 hrs after 48 hrs of fasting (lane 3). Total RNA was prepared from the above tissues using TRIzol. A real-time PCR method was used to quantify AdipoR mRNAs. The primer sets and probes were as described in the Experimental Procedures. The total large amount of each AdipoR transcript was obtained by normalization to β -actin transcript amount. The results are shown as the ratio to the AdipoR1 value in the liver of the unrestricted feeding group. Each bar represents the mean \pm standard error (S.E.) (n = 3) (*, p < 0.05; **, p < 0.01; compared with fasted mice).

Fig. 2 shows the levels of plasma insulin (A), blood glucose (B), and mRNA of AdipoR1 (C), and AdipoR2 (D) in control mice and STZ-treated mice, with or without insulin. Total RNA was prepared and AdipoR mRNAs were quantified as described in the legend of Fig. 1. The results are shown as the ratio to the AdipoR1 value in the skeletal muscle of non-STZ mice. Each bar represents the mean \pm S.E. (n = 3) (*, p < 0.05, **, p < 0.01, compared with STZ mice without insulin treatment).

Fig. 3 shows the amount of mRNAs of AdipoR1 (A, C, and E) and AdipoR2 (B, D, and F) in mouse hepatocytes (A and B) or C2C12 myocytes (C-F), treated with or without insulin, incubated with or without LY294002 or PD98059 (C and D) or transfected with adenoviruses containing LacZ or FoxoI (E and F). Hepatocytes were isolated from mouse liver and incubated as described in the Experimental Procedures. The cells were then incubated for 6 hrs with or without the indicated concentrations of insulin. Total RNA was prepared and AdipoR mRNAs were quantified as described in the legend of Fig. 1. The results are shown as the ratio to the AdipoR1 value in a control solvent. Each bar represents the mean \pm S.E. (n = 3) (*, p < 0.05, **, p < 0.01,

compared with cells treated with solvent).

Fig. 4 shows the expression of AdipoR1 (A) and AdipoR2 (B) mRNAs in liver, soleus muscle, EDL, WAT, and BAT, and the levels of blood glucose (C) and plasma insulin (D) in wild-type (C57BL/6) and ob/ob mice. Preparation of total RNA and
 5 quantification of AdipoR mRNAs were performed as described in the legend of Fig. 1. The results are shown as the ratio to the AdipoR1 value in the liver of wild-type mice. Each bar represents the mean \pm S.E. ($n = 3$)(*, $p < 0.05$, **, $p < 0.01$, between wild-type and ob/ob mice).

Fig. 5 shows the binding of globular (gAd) (A) or full-length adiponectin (Ad)
 10 (B) to the membrane fraction of skeletal muscles and AMPK activation by adiponectin in the skeletal muscle (C and D) of wild-type (C57BL/6) and ob/ob mice. Scatchard plot analysis of the binding of gAd or Ad to the membrane fraction of the skeletal muscle of wild type (C57BL/6) or ob/ob mice is shown (A and B). Phosphorylated α -subunit of AMPK in the skeletal muscle from wild type (C57BL/6) and ob/ob mice
 15 (C) was treated for 5 min with full-length adiponectin (Ad, 50 μ g body weight) or with solvent. Immunoblotting analysis was performed using anti-pAMPK and anti-AMPK antibodies. The amount of pAMPK was normalized by the amount of AMPK from the same sample. The results are shown as the ratio to the value for the solvent-treated wild-type mice. Each bar represents the mean \pm S.E. ($n = 5-7$)(*, $p < 0.05$, **, $p < 0.01$,
 20 compared with wild-type mice).

Best Mode for Carrying Out the Invention

The present invention relates to adiponectin receptor expression-enhancing agents comprising FoxoI protein or FoxoI gene. The present inventors discovered that
 25 insulin regulates the expression of adiponectin receptors, and that the regulation is mediated by the PI3-kinase signaling pathway, and confirmed that forced expression of FoxoI (constitutively active type), a member of the PI3-kinase signaling pathway, reduces the inhibition of adiponectin receptor (AdipoR1/R2) expression by insulin. Using the diabetes mouse model, the present inventors also discovered for the first time
 30 that AdipoR1/R2 expression levels regulate the binding of adiponectin and the activation of AMPK by adiponectin, referred to as "adiponectin sensitivity". FoxoI is a forkhead transcription factor, also termed FKHR. The FoxoI protein and FoxoI gene of the present invention include both the wild type and the constitutively active type. The amino acid sequence of mouse FoxoI (wild type) protein is shown in SEQ ID NO:
 35 1, and the cDNA sequence is shown in SEQ ID NO: 2. The amino acid sequence of the mouse FoxoI (constitutively active type) is shown in SEQ ID NO: 3. The amino

acid sequence of human FoxoI protein is shown in SEQ ID NO: 4, and the cDNA sequence is shown in SEQ ID NO: 5.

The FoxoI protein and FoxoI gene can be prepared by methods commonly known to those skilled in the art. For example, mRNAs prepared from cells
 5 expressing human or mouse FoxoI protein, such as skeletal muscle cells, hepatocytes, and adipose tissues, are used as templates to prepare a cDNA library according to standard methods. Hybridization is carried out using a part or the entire nucleotide sequence of SEQ ID NO: 2 or 5, or a part or the entire polynucleotide encoding the amino acid sequence of SEQ ID NO: 3 as a probe to prepare FoxoI cDNA.

10 Alternatively, the mRNAs prepared as above may be used as templates for RT-PCR with primers comprising a part of the nucleotide sequence of SEQ ID NO: 2 or 5, or a part of a polynucleotide encoding the amino acid sequence of SEQ ID NO: 3. Furthermore, the cDNA may be inserted into an appropriate expression vector, and introduced into host cells, expressed, and purified, to prepare a desired FoxoI protein.

15 The host and vector system may be selected depending on the purpose from an expression system such as a prokaryotic system, eukaryotic system, or cell free system, and a combination of vectors that match these expression systems. Examples of prokaryotic systems include *E. coli*, *Bacillus subtilis*, and *Actinomyces*. Yeasts, fungi, cultured animal cells, and such may be used as eukaryotic systems.

20 When *E. coli* is used as host, vectors such as pET, pPRO, and pBAD may be used. For efficient purification, vectors which express the desired protein as a fusion protein with other proteins may be selected. For example, vectors such as pGEX which express the desired protein as a fusion protein with GST (glutathione
 25 S-transferase), pCAL which expresses the desired protein as a fusion protein with CBP (calmodulin binding peptide), and pFLAG which expresses the desired protein as a fusion protein with an eight amino acid tag (Asp-Tyr-Lys-Asp-Asp-Asp-Lys) may be used. Examples of vectors for the *Bacillus subtilis* host include, pUB110, pAM α 1, and such, and an example of a vector for the *Actinomyces* host is pIJ702. Vectors such as YES2 and pESC may be used for yeasts. When using insect cells and mammalian
 30 cells as host, baculovirus may be used. An example of a baculovirus vector in mammalian cells is pAcCAGMCS1.

FoxoI proteins expressed in hosts can be purified by known techniques. FoxoI proteins expressed as a fusion protein with GST or His₆ may be purified by affinity chromatography on a glutathione sepharose column or nickel column,
 35 respectively. FoxoI proteins expressed using the above FLAG expression vector can be purified by affinity chromatography using commercially available monoclonal

antibodies against the FLAG peptide.

The FoxoI protein of the present invention includes "proteins structurally similar and functionally equivalent to the FoxoI protein" (described below as "FoxoI-like proteins") because proteins that are structurally similar to the above FoxoI protein and have a function similar to FoxoI protein can be also utilized as adiponectin receptor expression-enhancing agents. "FoxoI-like proteins" include naturally-occurring mutants of human or mouse FoxoI, FoxoI homologues and orthologues isolated from other organisms (mammals such as rats, guinea pigs, rabbits, dogs, cats, horses, sheep, donkeys, monkeys, chimpanzees, and birds), and artificial mutants of human or mouse FoxoI. More specifically, they are "proteins comprising an amino acid sequence with a deletion, substitution, addition and/or insertion of one or more amino acids in the amino acid sequence of SEQ ID NO: 1, 3, or 4", and "proteins encoded by a polynucleotide that hybridizes with the nucleotide sequence of SEQ ID NO: 2 or 5 under stringent conditions".

Herein, the number of amino acids that are added, deleted, inserted, and/or substituted is not limited as long as the action to reduce the insulin-mediated AdipoR1 and/or AdipoR2 expression inhibition is retained. Usually, the number of such mutations is 10% of the total number of amino acids or less, preferably 3% of the total number of amino acids or less, and more preferably 1% of the total number of amino acids or less.

FoxoI-like proteins can be prepared by methods commonly known to those skilled in the art. For example, the DNA of a protein with high homology to FoxoI may be isolated from skeletal muscle cells, adipocytes, and hepatocytes of mammals including humans under stringent hybridization conditions, using a part of the nucleotide sequence of SEQ ID NO: 2 or 5 as a probe, after which the FoxoI-like protein can be prepared by expressing the DNA. Alternatively, FoxoI-like proteins may be prepared by expressing a polynucleotide encoding a FoxoI-like protein, as described later.

Those skilled in the art can appropriately select the above-mentioned stringent hybridization conditions. For example, pre-hybridization is carried out in a hybridization solution containing 25% formamide (50% formamide under more stringent conditions), 4x SSC, 50 mM Hepes (pH 7.0), 10x Denhardt's solution, and 20 µg/ml denatured salmon sperm DNA at 42°C overnight. A labeled probe is added and hybridization is carried out by incubation at 42°C overnight. Post-hybridization washes are carried out under different levels of stringency including moderately stringent "1x SSC, 0.1% SDS, 37°C", highly stringent "0.5x SSC, 0.1% SDS, 42°C",

and more highly stringent "0.2x SSC, 0.1% SDS, 65°C" conditions. As the stringency level of post-hybridization washes increases, polynucleotides more highly homologous to the probe sequence are expected to be isolated. The above-described combinations of SSC, SDS, and temperature are merely examples of wash conditions. Those skilled
 5 in the art can achieve the same stringencies as those described above by appropriately combining the above factors or others (such as probe concentration, probe length, hybridization period, etc) that affect hybridization stringency.

A polypeptide encoded by the polynucleotide thus isolated using hybridization will usually comprise an amino acid sequence highly homologous to the polypeptides
 10 identified by the present inventors. "Highly homologous" refers to sequence homology of at least 40% or more, preferably 60% or more, further preferably 80% or more, further preferably 90% or more, further preferably at least 95% or more, further preferably at least 97% or more (for example, 98% to 99%). Amino acid sequence identity can be determined, for example, using the BLAST algorithm according to
 15 Karlin and Altschul (Proc. Natl. Acad. Sci. USA. 87:2264-2268, 1990; Proc. Natl. Acad. Sci. USA. 90: 5873-5877, 1993). A program designated BLASTX has been developed based on this algorithm (Altschul *et al.*, J. Mol. Biol. 215: 403-410, 1990). For analysis of amino acid sequence identity, the BLASTX program uses, for example, a score of 50 and a word length of 3 as parameters. When using BLAST and Gapped
 20 BLAST programs, the default parameters of the respective program are used. Specific methodology for performing BLAST analyses is publicly available at <http://www.ncbi.nlm.nih.gov>.

Whether the polypeptides prepared as above act to reduce insulin-mediated AdipoR1 and/or AdipoR2 expression inhibition may be confirmed by methods known to
 25 those skilled in the art. To give an example, the polypeptides may be expressed in mouse myogenic cell lines such as the C2C12 cell line, using adenoviruses as described in Example 6, the cells are then treated with insulin, and subsequently, the expression levels of AdipoR1/R2 mRNAs in the cell line may be measured using primers comprising the nucleotide sequence of SEQ ID NO: 10, 11, 13, or 14, to confirm the
 30 action to enhance AdipoR1 and/or AdipoR2 expression. Adenovirus vectors may be constructed according to the method described in Non-Patent Document 22.

In addition, polynucleotides encoding a FoxoI-like protein can be used, like FoxoI, as an adiponectin receptor expression-enhancing agent. The FoxoI gene of the present invention includes "polynucleotides encoding FoxoI-like proteins".
 35 Polynucleotides encoding FoxoI-like proteins of the present invention may be prepared by introducing site-specific or random mutations to the nucleotide sequence of SEQ ID

NO: 2 or 5, using genetic engineering methods commonly known to those skilled in the art, such as mutation introduction by PCR and cassette mutagenesis. Alternatively, the nucleotide sequence of SEQ ID NO: 2 or 5 introduced with a mutation(s) may be synthesized using a commercial DNA synthesizer.

5 The above FoxoI gene of the present invention can be used in gene therapy in humans and the like, by expressing FoxoI proteins and enhancing adiponectin receptor expression. Although the FoxoI gene may be used as is in the form of a polynucleotide as an adiponectin receptor expression-enhancing agent, it is preferable to
10 into an appropriate vector. The vector may be selected from vectors used in gene therapy, including for example retrovirus vectors, adenovirus vectors, adeno-associated virus (AAV) vectors, adenovirus/AAV hybrid vectors, artificial liposome vectors, etc.

 In addition, the present invention relates to adiponectin receptor expression-enhancing agents comprising a PI3-kinase/FoxoI pathway inhibitor. The
15 present inventors revealed for the first time that the regulation of adiponectin receptor expression by insulin is mediated by the PI3-kinase pathway. Examples of PI3-kinase-FoxoI pathway inhibitors include the PI3-kinase inhibitor LY294002 (CAS No.: [154447-36-6], 2-(4-morpholinyl)-8-phenyl-4H-1-benzopyran-4-one) and wortmannin (CAS No.: [19545-26-7]). A preferable example is LY294002, but the
20 inhibitors are not limited thereto. As long as a substance inhibits PI3-kinase activity, it is included in the adiponectin receptor expression-enhancing agents of the present invention. Moreover, other PI3-kinase-FoxoI pathway inhibitors such as Akt activity inhibitors and FoxoI phosphorylation inhibitors are also included in the present invention's adiponectin receptor expression-enhancing agents.

25 The present inventors showed for the first time that decreased adiponectin receptor expression leads to adiponectin resistance, and that insulin resistance is evoked as a result of impaired adiponectin actions. Therefore, the present invention's adiponectin receptor expression-enhancing agents can be used as agents for improving adiponectin resistance and insulin resistance. In addition to suppressing obesity,
30 diabetes, and arteriosclerosis by improving insulin resistance, adiponectin is also known to suppress arteriosclerosis through direct actions on arterial walls. Thus, the adiponectin receptor expression-enhancing agents of the present invention could be very useful as obesity-preventing or improving agents, diabetes-preventing or treating agents, or arteriosclerosis-preventing or treating agents.

35 When using a FoxoI protein, a FoxoI gene, or a PI3-kinase-Akt pathway inhibitor of the present invention as a pharmaceutical agent, the substance can either be

directly administered to a patient, or administered as a formulation prepared by a commonly known pharmaceutical method. For example, the substance may be appropriately combined with a pharmaceutically acceptable carrier or vehicle, more specifically, sterilized water, saline, plant oil, emulsifying agent, suspension, detergent, stabilizing agent, or such for formulation, and then administered. Administration to patients may be carried out by known methods such as intraarterial administration, intravenous administration, subcutaneous administration, intramuscular administration, oral administration, and transdermal administration. The dosage varies depending on the age and body weight of patients and administration method, but those skilled in the art can appropriately select a suitable dose.

Furthermore, the present invention relates to adiponectin receptor expression-suppressing agents comprising insulin or insulin gene. As described above, insulin was shown to downregulate adiponectin receptor expression and thus evokes adiponectin resistance and insulin resistance. The adiponectin receptor expression-suppressing agents comprising insulin or insulin gene are important as reagents for further elucidating the mechanism of adiponectin resistance and insulin resistance. The amino acid sequence of mouse insulin is shown in SEQ ID NO: 6 and the cDNA sequence of mouse insulin is shown in SEQ ID NO: 7. The amino acid sequence of human insulin is shown in SEQ ID NO: 8 and the cDNA sequence of human insulin is shown in SEQ ID NO: 9.

Insulin and insulin gene can be prepared by methods commonly known to those skilled in the art. For example, insulin gene may be obtained by preparing mRNA from mouse or human pancreatic cells, and performing RT-PCR using a part of the nucleotide sequence of SEQ ID NO: 7 or 9 as primer. Alternatively, a cDNA library prepared from mouse or human pancreatic cells may be screened using a part of the nucleotide sequence of SEQ ID NO: 7 or 9 as a probe to prepare the insulin gene. An insulin gene prepared as above may be expressed using an appropriate host-vector system for purification to prepare a desired insulin protein.

The above insulin protein is thought to be able to reduce the amount of adiponectin receptors expressed in cells by contacting it with cells that express adiponectin receptors. Similarly, an insulin gene introduced into adiponectin receptor-expressing cells can produce cells with reduced adiponectin receptor expression amount. Such cells may be used for elucidating the mechanisms of adiponectin resistance, insulin resistance, and diabetes.

Furthermore, the insulin gene may be introduced into animals such as mice, to produce animals with sustainably reduced adiponectin receptor expression levels. The

insulin gene may be introduced into animals by introducing a fragment of the insulin gene by the particle gun method and such, but preferably, it is introduced using the aforementioned vectors for gene therapy. The above animals are thought to be useful as animal models for adiponectin resistance, insulin resistance, and diabetes, and especially for elucidating the mechanisms and for developing therapeutic agents and therapeutic methods.

Furthermore, the present invention relates to methods of screening for adiponectin receptor expression-enhancing agents, adiponectin resistance-improving agents, insulin resistance-improving agents, obesity-improving agents, or for therapeutic agents for diabetes or arteriosclerosis, in which the methods of screening comprise the step of contacting test substances with PI3 kinase and PI3 kinase substrates. As described in the Examples, a PI3-kinase inhibitor was confirmed to inhibit the suppression of adiponectin receptor expression by insulin, clarifying that the inhibition of adiponectin receptor expression by insulin is mediated through the PI3-kinase pathway. Therefore, the activity to inhibit PI3-kinase can be used as a marker for selecting adiponectin receptor expression-enhancing agents.

The first embodiment of the present invention comprises an *in vitro* system using purified PI3-kinase. The above system comprises the steps of contacting a test substance with PI3-kinase and a PI3-kinase substrate, detecting the PI3-kinase activity in the presence of the test substance, comparing the activity of PI3-kinase with or without the test substance, and selecting a test substance when PI3-kinase activity in the presence of the test substance is lower than that in the absence of the test substance. PI3-kinase catalyzes the reaction that produces $PI(3,4,5)P_3$ using $PI(4,5)P_2$ as a substrate. The above reaction activity can be measured by known methods. For example, phosphorylation of phosphatidylinositol substrates by PI3-kinase may be carried out with P^{32} or P^{33} in the presence or absence of a test substance. Radioactive phosphorylation products are extracted, and then separated and detected by thin-layer chromatography. By comparing the activities of PI3-kinase with or without the test substance, the test substance's inhibition of PI3-kinase activity can be measured. In addition, PI3-kinase may be reacted with a substrate in the presence of a test substance, and the thus produced $PI(3,4,5)P_3$ may be measured by ELISA. This method can be conducted easily using a commercial PI3-kinase ELISA kit (Echelon Biosciences, Inc). In addition, a FRET (fluorescence resonance energy transfer)-based PI3-kinase assay (PI Profiler™, Upstate) may be used for the measurement.

The second embodiment of the present invention includes a system which comprises the step of contacting a test substance with PI3-kinase and a PI3-kinase

substrate in cells, by incubating PI3-kinase-expressing cells with the test substance. In this system, following the step of contacting the test substance with PI3-kinase in the cells, the following steps can be carried out: measuring PI(3,4,5)P₃ in the cells; comparing the PI(3,4,5)P₃ measurements in the absence or presence of the test substance; and selecting the substance when the measured value in the presence of the test substance is lower than that in the absence of test substance. PI(3,4,5)P₃ may be measured by preparing cell lysates, and PI(3,4,5)P₃ in the cell lysates may be measured by an immunoassay using a specific antibody.

Examples of test substances according to the present invention include nucleic acids, proteins, polypeptides, synthetic low molecular weight compounds, cell extracts, and extracts of natural products, but are not restricted thereto, and may be other substances. Test substances obtained by a screening method of the present invention are expected to enhance adiponectin receptor expression through PI3-kinase inhibition, and thus may be used as adiponectin receptor expression-enhancing agents, adiponectin resistance-improving agents, insulin resistance-improving agents, obesity-improving agents, or therapeutic agents for diabetes or arteriosclerosis.

Moreover, the present invention relates to methods of screening for adiponectin receptor expression-enhancing agents, adiponectin resistance-improving agents, insulin resistance-improving agents, obesity-improving agents, or therapeutic agents for diabetes or arteriosclerosis, in which the screening methods comprise the step of contacting Akt with FoxoI. In the methods of the present invention, phosphorylation of FoxoI is used as a marker for screening the above substances. The Examples of the present invention show that insulin inhibits AdipoR1/R2 expressions via inactivation of FoxoI. Accordingly, phosphorylation of FoxoI at specific residues suppresses the activity of FoxoI to transcribe target genes. On the other hand, the promoter region of AdipoR1/R2 contains a cis-acting element that is capable of binding to FoxoI. These suggest that FoxoI is phosphorylated by Akt and inactivated, which in turn suppresses adiponectin receptor expression. Thus, phosphorylated FoxoI measurement can be used as a marker when screening for adiponectin receptor expression-enhancing agent.

As one embodiment of the present invention, a method of the present invention may be performed as an *in vitro* assay system, using purified Akt and FoxoI. The above system comprises the steps of contacting a test substance with Akt and FoxoI, measuring the phosphorylated FoxoI in the presence or absence of the test substance, comparing the measurements of phosphorylated FoxoI with or without the test substance, and selecting the test substance when the measured value of phosphorylated FoxoI in the presence of the test substance is less than the activity in the absence of the

test substance. The above method can be performed using known methods. For example, Akt is contacted with FoxoI in the presence of a test substance *in vitro*, and then reacted with phospho-specific antibodies to measure the amount of phosphorylated FoxoI by an immunoassay. The phospho-specific antibody preferably detects residues phosphorylated by Akt. Examples of such an antibody include the commercially available Phospho-FKHR(Ser256) Antibody, and Phospho-FKHR(Thr24)/FKHR1(Thr32) Antibody (both from Cell Signaling Technology).

Another embodiment of the methods of the present invention includes a system comprising the step of contacting a test substance with Akt and FoxoI in cells, by incubating Akt and FoxoI expressing cells in the presence or absence of the test substance. In this system, following the above step, the following steps are performed: measuring the phosphorylated FoxoI in the cells; comparing the measurements of phosphorylated FoxoI in the presence or absence of the test substance; and selecting the test substance when the measured value of phosphorylated FoxoI in the presence of the substance is less than the activity in the absence of the test substance. The phosphorylated FoxoI in the cells may be measured, for example, by preparing cell lysates, and then measuring by a method similar to that described above.

Examples of test substances for the methods of the present invention include nucleic acids, proteins, polypeptides, synthetic low molecular weight compounds, cell extracts, and extracts of natural products, but are not restricted thereto, and may be other substances. Substances obtained by the screening methods of the present invention are expected to have an action to suppress the inactivation of FoxoI by insulin, and thereby enhance adiponectin receptor expression. Thus, such substances may be used as adiponectin receptor expression-enhancing agents, adiponectin resistance-improving agents, insulin resistance-improving agents, obesity-improving agents, or therapeutic agents for diabetes or arteriosclerosis.

All prior art references cited herein are incorporated into this description by references.

Examples

The present invention will be explained more specifically below based on Examples; however, the present invention is not limited thereto. All the reagents used in the Examples were obtained from the sources described in references (Non-Patent Documents 6, 10, 14, and 19).

[Example 1] Fasting-induced expression increase and refeeding-induced expression decrease of AdipoR1/R2 genes

To elucidate whether or not the expression of AdipoR1 and/or AdipoR2 is regulated under physiological and/or pathophysiological conditions, the expression levels of AdipoR1 and AdipoR2 in insulin and adiponectin target organs were examined in mice under conditions of unrestricted feeding, fasting, and refeeding.

Fifteen-week male C57BL/6 mice obtained from Charles River Breeding Laboratories (Washington, MA) were kept in colony cages, maintaining a 12-hr light/dark cycle. The mice were provided or deprived of food for 48 hrs from the start of the dark cycle (9:00 p.m.). The refed group was provided with food after 48 hrs of fasting, and then sacrificed 6 hrs after refeeding for tissue isolation. The mice were given a high fat diet containing 1152 g of fat (Benibana (safflower oil, high oleic acid type), Japan, containing 46% oleic acid (18:1n-9) and 45% linolenic acid (18:2n-6) in total fatty acids), 1191.6 g of casein (No. 19, Oriental Yeast, Tokyo, Japan), 633.6 g of sucrose (No. 13, Oriental Yeast, Tokyo), 50.4 g of a vitamin mix (No. 20 (AIN76), Oriental Yeast), 352.8 g of a mineral mix (No. 20 (AIN76), Oriental Yeast), 201.6 g of cellulose powder (No. 19, Oriental Yeast), 18 g of DL-methionine (Wako Pure Chemical Industries, Osaka, Japan), and 360 ml of water; 3600 g in total (Non-Patent Document 19).

Real time PCR was performed to quantify Adipo receptor mRNAs (Non-Patent Document 14). Total RNA was prepared from cells or tissues using TRIzol (GIBCO/BRL) according to the instructions provided by the manufacturer. The relative amount of each AdipoR transcript was obtained by normalization to the amount of actin transcript in the same cDNA (Non-Patent Document 14). The primer sets and probes used for the quantification are shown below.

mAdipoR1

Forward primer:	acgttgagagtcacccgtat	(SEQ ID NO: 10)
Reverse primer:	ctctgtgtggatgcggaagat	(SEQ ID NO: 11)
Probe:	cctgctacatggccacagaccacct	(SEQ ID NO: 12)

mAdipoR2

Forward primer:	tcccaggaagatgaagggttat	(SEQ ID NO: 13)
Reverse primer:	ttccattcgttcgatagcatga	(SEQ ID NO: 14)
Probe:	atgtccccgctcctacaggccc	(SEQ ID NO: 15)

The expression levels of AdipoR1 and AdipoR2 mRNAs in the liver

dramatically increased after 48 hrs of fasting, and were rapidly restored by refeeding to the same level as in natural feeding conditions (Figs. 1A and 1B). A similar AdipoR2 mRNA expression regulation due to fasting and refeeding was also observed in skeletal muscles (Figs. 1C and 1D).

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[Example 2] An inverse correlation between the plasma insulin level and the expression level of AdipoR1/R2 gene

The above results in Example 1 confirm that the expression of AdipoR1/R2 gene is regulated in response to fasting and feeding, suggesting the involvement of a particular hormone(s) or nutrient(s) in the above regulation. Because insulin is a classic nutritional hormone, plasma insulin levels of the above mice under unrestricted-feeding, fasting, and refeeding states were measured using an insulin immunoassay (Shibayagi, Gunma, Japan).

As expected, fasting decreased the plasma insulin level (Fig. 1F), whereas refeeding increased the plasma insulin level in parallel with the blood glucose level (Fig. 1E). These findings suggest that the plasma insulin values inversely correlate with the expressed amount of AdipoR1/R2 mRNAs, presenting the possibility that insulin may control AdipoR1/R2 expression.

20 [Example 3] STZ treatment-induced expression increase and insulin-induced expression decrease of AdipoR1/R2

To further examine the effect of insulin on AdipoR1/R2 expression, mice were treated with streptozotocin (STZ) and AdipoR1/R2 expression was observed. STZ disrupts pancreatic β cells, and causes acute insulin deficiency. To induce diabetes, mice were intraperitoneally administered twice with 0.2-0.3 ml of a 50 mM sodium citrate solution (pH 4.5) containing STZ (200 mg/kg body weight). Control mice were injected with a 50 mM sodium citrate solution (pH 4.5). For three days after injection, plasma glucose levels were measured, and diabetes was confirmed (glucose value > 250 mg/dl). Human regular insulin (1 unit/kg, Eli Lilly, Indianapolis, IN) dissolved in 0.2 ml of saline was administered to the STZ + insulin treated group. The mice in the STZ + saline treated group and non-STZ treated group were intraperitoneally administered with 0.2 ml of saline. After insulin or saline administration, the animals were fasted for 6 hrs and then sacrificed to collect the hind limb skeletal muscles.

STZ treatment made plasma insulin disappear (Fig. 2A), and at the same time, caused a significant increase of plasma glucose (Fig. 2B, lanes 1 and 2). The STZ and insulin treated group exhibited a lower plasma glucose value as compared to the group

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treated with STZ only (Fig. 2B, lanes 2 and 3).

Next, the amount of AdipoR1 and AdipoR2 mRNA in skeletal muscle and liver extracts was measured as described above. In STZ-treated mice, the AdipoR1 and AdipoR2 mRNA in the skeletal muscle respectively showed a 103% and 38% increase (Figs. 2C and 2D), which was almost completely restored by insulin treatment. These findings further support the possibility that insulin reduces AdipoR1/R2 mRNA levels. STZ treatment had little effect on the AdipoR1/R2 mRNA levels in the liver, whereas insulin significantly reduced the AdipoR1/R2 mRNA levels in the liver of STZ-treated mice (data not shown). These data may be explained by the findings that inflammation may decrease AdipoR1/R2 mRNA levels (Tsuchida, Yamauchi, and Kadowaki, manuscript in preparation) and that STZ treatment can induce inflammation in the liver (Non-Patent Document 23).

[Example 4] Decrease in *in vitro* AdipoR1/R2 mRNA levels due to insulin.

The direct effect of insulin on AdipoR1/R2 expression in Hepatocytes was examined. Hepatocytes were prepared from fasted 8-week-old male C57BL/6 mice by collagenase perfusion method (Non-Patent Document 20). Aliquots of 0.8×10^6 cells were plated into Williams medium E supplemented with 5% (v/v) fetal calf serum, 10 nM dexamethasone, 1 nM insulin, 100 units/ml penicillin, and 100 μ g/ml streptomycin, on a collagen I-coated 12-well dish. After incubation at 37°C, 5% CO₂ for 5 hrs, the cells were washed twice with PBS, and then incubated for 6 hrs in Williams medium E supplemented with 1% bovine serum albumin. The cells were then transferred to Williams medium E supplemented with 1% bovine serum albumin and 100 nM insulin, and incubated with or without 100 nM insulin for 6 hrs. The hepatocytes were harvested to measure the total amount of AdipoR1/R2 mRNA. Insulin reduced the total amount of AdipoR1/R2 mRNA (Figs. 3A and 3B).

To further explore the possibility that insulin may reduce AdipoR1/R2 mRNA levels, the effect of insulin on the expression levels of AdipoR1/R2 mRNA in C2C12 myocytes was observed *in vitro*. Induction of myogenic differentiation was carried out according to a previously reported method (Non-Patent Document 10). Briefly, mouse C2C12 myoblasts were cultured in Dulbecco's modified Eagle's medium- high glucose- (DMEMH; 90%) with 10% (v/v) FBS. When the cells reached 80% confluence, the medium was replaced with a low serum differentiation medium (98% DMEMH, 2.5% (v/v) horse serum). The medium was changed every day to induce the differentiation of myoblasts into myotubes. By day 5, the cells had differentiated into multinucleated contractile myotubes. Treatment with 10 nM insulin indeed reduced the expression

levels of AdipoR1/R2 mRNA in the above C2C12 myocytes after 6 hrs (Figs. 3C and 3D).

[Example 5] Suppression of insulin-induced reduction in AdipoR1/R2 mRNA levels by a phosphoinositide 3-kinase (PI3-kinase) inhibitor

C2C12 cells were incubated in DMEMH supplemented with 1% bovine serum albumin for 12 hrs, and then exposed to insulin transduction pathway inhibitors (10 μ M LY294002 or 10 μ M PD98059) 30 min before the addition of insulin. The cells were harvested 6 hrs after the addition of insulin to measure the AdipoR1/R2 mRNA levels in the C2C12 myocytes treated with insulin for 6 hrs (Figs. 3C and 3D). The PI3-kinase inhibitor LY294002 canceled the AdipoR1/R2 mRNA level reduction effect of insulin. In contrast, PD98059, an inhibitor of mitogen-activated protein kinase activation, did not essentially affect the response to insulin. The above results led to the conclusion that the induction of AdipoR1/R2 gene expression may be mediated by the PI3-kinase branched pathway of the insulin signal transduction pathway. The inhibitory effects shown in Figs. 3C and 3D exhibited maximum value against all of the reagents tested. Further, the points were verified by dose-response experiments (data not shown).

Many, if not all, of the known actions of insulin are ascribable to the tyrosine phosphorylation in which the insulin receptor of IRS-1 and IRS-2 plays a part, and this phosphorylation triggers kinase cascades involving the PI3-kinase/Akt pathway and the MAP kinase pathway (Non-Patent Documents 26-28). The effect of insulin on AdipoR1/R2 gene expression was hardly affected by the MEK inhibitor PD98059, and thus does not appear to require activation of the mitogen-activated protein kinase cascade. In contrast, the results with LY294002 clearly implicated the PI3-kinase signal transduction pathway. The results with the PI3-kinase inhibitor leads to the conclusion that the induction of AdipoR1/R2 gene expression may be mediated by the PI3-kinase branched pathway of the insulin signal transduction pathway.

[Example 6] Enhancement of AdipoR1/R2 mRNA levels by FoxoI and inhibition of insulin-induced reduction of AdipoR1/R2 mRNA levels

The PI3-kinase signal transduction pathway is thought to be involved in the insulin-mediated regulation of several genes, particularly, the regulation of the gene for insulin-like growth factor binding protein (IGF-BP1) (Non-Patent Document 29). Recently, Akt, an effector downstream of PI3-kinase, was shown to phosphorylate several transcriptional activators of the forkhead family including FoxoI *in vitro* and in intact cells. When specific residues of these activators are phosphorylated, the

phosphorylated activators become isolated in the cytoplasm, and as a result, their ability to activate the transcription of target genes is inhibited (Non-Patent Documents 30-32). Because IGF-BP1 gene contains in its promoter a *cis*-acting element capable of binding to FoxoI, the insulin dependent suppression of IGF-BP1 gene could, at least partially, be due to Akt-mediated phosphorylation and inactivation of the forkhead transcriptional
 5 activators (Non-Patent Document 29).

The AdipoR1/R2 genes also contain in their promoters a *cis*-acting element capable of binding to FoxoI. Thus, the insulin dependent suppression of AdipoR1/R2 genes could, at least partially, be due to PI3-kinase/Akt-mediated phosphorylation and
 10 inactivation of the forkhead transactivators (Non-Patent Documents 21 and 22). Thus, the effect of FoxoI expression on insulin dependent suppression of AdipoR1/R2 genes was observed. C2C12 cells were transfected with adenoviruses containing wild-type mouse FoxoI cDNA and constitutively active form of FoxoI (T24A/S253D/S316A (ADA)) cDNA (Non-Patent Documents 21 and 22) according to a previously described
 15 method (Non-Patent Document 10) with minor modifications. Differentiation was induced as described above. At day 5, the above cells were treated with the indicated concentrations of insulin, and then analyzed.

The expression of constitutively active FoxoI (FoxoI-ADA) induced a dramatic increase in AdipoR1/R2 mRNA levels. Insulin failed to restore the above effect of
 20 FoxoI (Figs. 3E and 3F). These findings suggest that FoxoI enhances AdipoR1/R2 expression levels and that insulin suppresses AdipoR1/R2 mRNA expressions through inactivation of FoxoI.

[Example 7] Downregulation of AdipoR1/R2 expression levels in ob/ob mice

25 The expression of AdipoR1 and R2 in insulin-sensitive tissues was examined using insulin-resistant ob/ob mice and C57BL/6 mice as a control. The ob/ob mice are genetically obese mice that exhibit obesity, fatty liver, and hyperinsulinemia, due to a deficiency in the leptin gene. Insulin-sensitive tissues such as liver, skeletal muscles (soleus muscle and extensor digitorum longus (EDL)), and adipose tissues (white
 30 adipose tissue (WAT) and brown adipose tissue (BAT)) were examined. The results are shown in Figs. 4A and 4B. In the control C57BL/6 mice, AdipoR1 was expressed most abundantly in skeletal muscles, whereas AdipoR2 was expressed most in the liver. Interestingly, AdipoR1 and AdipoR2 were both expressed more abundantly in EDL than in soleus muscles. Both AdipoR1 and R2 were expressed in both WAT and BAT
 35 adipose tissues.

The expressions of both AdipoR1 and R2 in ob/ob mice were significantly less

than in C57BL/6 mice in all of the examined tissues except the liver (Figs. 4A and 4B). The expression level of AdipoR1 in the liver, soleus muscle, EDL, WAT, and BAT of the ob/ob mice was 79.4%, 54.5%, 50.5%, 31.9%, and 28.1% of the level in the wild-type mice, respectively (Fig. 4A). The expression level of AdipoR2 in the liver, soleus muscle, EDL, WAT, and BAT of the ob/ob mice was 63.3%, 51.4%, 59.5%, 7.9%, and 29.4% of that in the wild-type mice, respectively (Fig. 4B).

[Example 8] The correlation between reduction in AdipoR1/R2 expression levels and "adiponectin sensitivity" in ob/ob mice

10 The association between AdipoR1/R2 expression levels and the effects of adiponectin in each tissue was examined. Adiponectin contributes to the enhancement of insulin-sensitivity via activation of AMPK and PPAR α . Thus, the activation of AMPK by adiponectin and the amount of adiponectin bound to the membrane fraction of skeletal muscles were measured in obesity-related insulin resistance model mice
15 (ob/ob mice) and wild-type mice (C57BL/6 mice).

 The amount of adiponectin bound to the membrane fraction of skeletal muscles was determined as follows. First, recombinant mouse full-length adiponectin (Ad) was prepared as described in references (Non-Patent Documents 10 and 14). Tyrosine of synthetic adiponectin was labeled with ^{125}I using IODO-beads (Pierce) in the presence
20 of Na ^{125}I (2,000 Ci/mmol, Amersham Pharmacia Biotech) according to the manufacturer's protocol. The membrane fraction of skeletal muscles was purified by sucrose gradient centrifugation, and incubated at 4°C for 1 hr in a binding buffer (ice-cold phosphate-buffered saline/10% skim milk) containing an indicated concentration of [^{125}I]-adiponectin (2,000 cpm/ng protein) and non-labeled competitors.
25 Two hours later, when binding experiments were conducted at 4°C, establishment of binding equilibrium was confirmed. The cells were washed three times with ice-cold phosphate-buffered saline, and the membrane-bound radioactivity was determined using a γ -counter (Non-Patent Documents 10 and 14). Non-specific binding was determined by over 200-fold unlabeled adiponectin. Specific binding was calculated by
30 subtracting the nonspecific binding from the total binding. The values shown in the results are the average of three determinations from three experiments.

 The activation of AMPK by adiponectin was measured as follows. Fifteen-week-old male ob/ob mice or age-matched C57BL/6 mice were anesthetized with pentobarbital, and then injected with the above recombinant adiponectin (50 $\mu\text{g}/10$
35 g of body weight) through the inferior vena cava. Skeletal muscles were collected five minutes after adiponectin injection. The amount of phosphorylated AMPK in skeletal

muscles was determined by western blotting (Non-Patent Documents 10 and 14).

Interestingly, the binding of adiponectin was more robust in the membrane fraction of skeletal muscles of wild-type mice than that of ob/ob mice (Figs. 5A and 5B). Scatchard plot analysis shows that both the high affinity binding site and low affinity
 5 binding site for globular adiponectin and full-length adiponectin were decreased in the skeletal muscles of ob/ob mice compared with those in wild-type mice, suggesting that the numbers of both AdipoR1 and AdipoR2 were decreased (Figs. 5A and 5B) (Non-Patent Document 14). Moreover, adiponectin could activate AMPK in the skeletal muscles of wild-type mice, but could not do so in the skeletal muscles of ob/ob
 10 mice (Figs. 5c and 5D). These findings suggest that AdipoR1/R2 expression levels regulate adiponectin binding and activation of AMPK by adiponectin, referred to as "adiponectin sensitivity".

Thus, the present inventors discovered that AdipoR1/R2 expressions are downregulated in the obesity-related insulin-resistance model ob/ob mice. Previously,
 15 the inventors showed that plasma adiponectin level was reduced in ob/ob mice (Non-Patent Document 12), and that such changes in plasma adiponectin level play a causal role in the regulation of insulin sensitivity. Taken together, these findings raise an interesting possibility that changes in AdipoR1/R2 expression levels, in addition to the plasma adiponectin level, may play a causal role in the regulation of insulin
 20 sensitivity. This possibility is consistent with the previous report by the inventors that the AdipoR1 or AdipoR2 expression reduction by their corresponding siRNA is associated with the insulin-sensitizing effect of adiponectin (Non-Patent Document 14). The present inventors proved for the first time that AdipoR1 or AdipoR2 expression reduction in model mice is associated with the insulin-sensitizing effect of adiponectin
 25 *in vivo*.

It is expected that chronic overeating and its consequent increase in insulin level will lead to decreased AdipoR1/R2 expressions. Such a decrease will cause both adiponectin resistance and insulin resistance. The decrease in AdipoR1/R2 mRNAs leads to a decrease in adiponectin binding, which in turn leads to a decrease in the
 30 adiponectin effects, referred to as "adiponectin resistance", completing a so-called "vicious cycle". These findings may provide a novel molecular mechanism against long-term insulin elevation that leads to insulin resistance, comprising downregulation of adiponectin receptors in addition to the well known downregulation of insulin receptors.

35 To conclude, AdipoR1/R2 expressions in insulin target tissues appear to be inversely correlated with the plasma insulin level. The present invention demonstrated

for the first time that insulin negatively regulates the expression level of adiponectin receptors through the PI3-kinase/FoxoI pathway. These data suggest that not only AdipoR1/R2 agonists but also a strategy to increase AdipoR1/R2 can be a logical approach to provide a novel treatment concept for insulin resistance and type 2 diabetes.

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Industrial Applicability

Adiponectin receptor expression-enhancing agents of the present invention are capable of resolving both adiponectin resistance and insulin resistance, and thus will be useful as adiponectin resistance-improving agents, insulin resistance-improving agents, 10 obesity-improving agents, or therapeutic agents for diabetes or arteriosclerosis. The screening methods of the present invention makes use of the fact that insulin regulates adiponectin receptor expression via the PI3-kinase pathway, and will be particularly useful for discovering candidate substances for adiponectin resistance-improving agents, insulin resistance-improving agents, obesity-improving agents, or therapeutic agents for 15 diabetes or arteriosclerosis.